IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Group Art Unit: Not yet assigned

Ashkenazi et al.

Examiner: Not yet assigned

Serial No.: Not yet assigned

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For: Secreted and Transmembrane

Polypeptides and Nucleic Acids

Encoding the Same

PRELIMINARY AMENDMENT

Assistant Commissioner of Patents

Washington, D.C. 20231

sir:

Prior to substantive examination of the above captioned patent application (which is filed herewith), and for calculation of the proper filing fee, Applicants respectfully request that the following amendments be entered.

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In the claims:

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Please cancel Claims 1-57 without prejudice or disclaimer.

Please add new Claims 58-77 as follows.

- -58.(New) An isolated nucleic acid having at least 80% nucleic acid sequence identity to:
- a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID (a) NO:206);
- a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID (b) NO:206), lacking its associated signal peptide;
- (c) a nucleic acid seq in Figure 75 (SEQ ID NO:206); a nucleic acid sequence encoding the extracellular domain of the polypeptide shown
 - a nucleic acid sequence encoding the extracellular domain of the polypeptide shown (d) in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;
 - the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205); (e)
 - (f) the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEO ID NO:205); or
- the full-length coding sequence of the cDNA deposited under ATCC accession (g) number 209812.
 - (New) The isolated nucleic acid of Claim 58 having at least 85% nucleic acid 59. sequence identity to:
 - a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID (a) NO:206);
 - (b) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal pertide:

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(c) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206);

- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;
 - (e) the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205);
- (f) the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205); or
- (g) the full-length coding sequence of the cDNA deposited under ATCC accession number 209812.
- 60. (New) The isolated nucleic acid of Claim 58 having at least 90% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID NO:206);
 - (b) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;
 - (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206);
 - (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;
 - (e) the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205);
 - (f) the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205); or
 - (g) the full-length coding sequence of the cDNA deposited under ATCC accession number 209812.

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61. (New) The isolated nucleic acid of Claim 58 having at least 95% nucleic acid sequence identity to:

- a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID (a) NO:206);
- a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID (b) NO:206), lacking its associated signal peptide;
- a nucleic acid sequence encoding the extracellular domain of the polypeptide shown (c) in Figure 75 (SEQ ID NO:206);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown [=
 - the nucleic acid sequence shown in Figure 74 (SEO ID NO:205):
- in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;

 (e) the nucleic acid sequence shown in Figure 74 (SEQ ID

 (f) the full-length coding sequence of the nucleic acid sequen the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEQ
- (g) the full-length coding sequence of the cDNA deposited under ATCC accession (g) the state of t
 - (New) The isolated nucleic acid of Claim 58 having at least 99% nucleic acid sequence identity to:
 - (a) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEO ID NO:206);
 - a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID (b) NO:206), lacking its associated signal peptide;
 - a nucleic acid sequence encoding the extracellular domain of the polypeptide shown (c) in Figure 75 (SEQ ID NO:206);
 - a nucleic acid sequence encoding the extracellular domain of the polypeptide shown (d) in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;

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- the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205); (e)
- (f) the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205); or
- the full-length coding sequence of the cDNA deposited under ATCC accession (g) number 209812.
 - 63. (New) An isolated nucleic acid comprising:
- (a) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID NO:206);
- (b) a nucleic acid sequence encoding NO:206), lacking its associated signal peptide; a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID
- a nucleic acid sequence encoding the extracellular domain of the polypeptide shown (c) in Figure 75 (SEQ ID NO:206);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;

 (e) the nucleic acid sequence shown in Figure 74 (SEQ ID the full-length coding sequence of the nucleic acid sequence.
 - the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205);
 - the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205); or
 - (g) the full-length coding sequence of the cDNA deposited under ATCC accession number 209812.
 - (New) The isolated nucleic acid of Claim 63 comprising a nucleic acid sequence 64. encoding the polypeptide shown in Figure 75 (SEQ ID NO:206).

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65. (New) The isolated nucleic acid of Claim 63 comprising a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide.

- 66. (New) The isolated nucleic acid of Claim 63 comprising a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206).
- (New) The isolated nucleic acid of Claim 63 comprising a nucleic acid sequence 67. encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking
- its associated signal peptide.

 68. (New) The isomorphisms shown in Figure 74 (SEQ ID (New) The isolated nucleic acid of Claim 63 comprising the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205).
 - 69. (New) The isolated nucleic acid of Claim 63 comprising the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205).
- 70. (New) The isolated nucleic acid of Claim 63 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 209812.
 - 71. (New) An isolated nucleic acid that hybridizes to:
 - (a) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEO ID NO:206);
 - (b) a nucleic acid sequence encoding the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;

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(c) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206);

- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 75 (SEQ ID NO:206), lacking its associated signal peptide;
 - (e) the nucleic acid sequence shown in Figure 74 (SEQ ID NO:205);
- (f) the full-length coding sequence of the nucleic acid sequence shown in Figure 74 (SEO ID NO:205); or
- (g) the full-length coding sequence of the cDNA deposited under ATCC accession number 209812.
- 72. (New) The isolated nucleic acid of Claim 71, wherein said hybridization occurs under stringent conditions.
- SOLZCES ... CELOI 73. (New) The isolated nucleic acid of Claim 71 which is at least 10 nucleotides in length.
 - 74. (New) A vector comprising the nucleic acid of Claim 58.
 - (New) The vector of Claim 74, wherein said nucleic acid is operably linked to control 75. sequences recognized by a host cell transformed with the vector.
 - 76. (New) A host cell comprising the vector of Claim 74.
 - 77. (New) The host cell of Claim 76, wherein said cell is a CHO cell, an E. coli or a yeast cell.--

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Applicants respectfully request entry of these new claims for prosecution in this application. The Examiner is invited to contact the undersigned at (650) 225-4563 if any issues may be resolved in that manner.

Respectfully submitted,

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